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GenCore version 5.1.6
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OM nucleic - nucleic search, using sw model

Run on: June 9, 2003, 03:30:13 ; Search time 425 Seconds
(without alignments)
3734.295 Million cell updates/sec

Title: US-10-091-628-1

Perfect score: 1134

Sequence: 1 atgagagcgaatgcttcacg.....acatcattcatgtaag 1134

Scoring table:

Gapop 10.0, Gapext 1.0

Reached: 870385 segs, 699768693 residues

Total number of hits satisfying chosen parameters: 1740770

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

Published Applications NA:
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2: /cgn2_6/ptodata/1/pubpna/PTC_NEW_PUB.seq:*
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14: /cgn2_6/ptodata/1/pubpna/US60_PUBCOMB.seq:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length DB	ID	Description
1	1134	100.0	1134	9 US-10-091-628-1	Sequence 1, Appl1
2	1134	100.0	1600	9 US-10-091-628-3	Sequence 3, Appl1
3	183.2	16.2	1663	10 US-09-917-800A-1626	Sequence 1626, Ap
4	173.6	15.3	1580	10 US-09-880-107-2176	Sequence 2176, Ap
5	79.8	7.0	360	10 US-09-864-761-31375	Sequence 31375, A
6	79.8	7.0	560	10 US-09-864-761-14847	Sequence 14847, A
7	77.8	6.9	310	10 US-09-833-381-317	Sequence 317, App
8	74	6.5	401	10 US-09-960-352-2253	Sequence 2253, Ap
9	67.6	6.0	972	9 US-09-738-626-2554	Sequence 2554, Ap
10	67.6	6.0	3309400	9 US-09-738-626-1	Sequence 1, Appl1
11	64.4	5.7	1017	9 US-09-938-842A-380	Sequence 380, App
12	62.4	5.5	1317	9 US-10-091-628-4	Sequence 4, Appl1
13	62.4	5.5	1777	9 US-10-091-628-6	Sequence 6, Appl1
14	58.6	5.2	374	10 US-09-833-381-318	Sequence 318, App
15	54.2	4.8	407	10 US-09-960-352-10081	Sequence 10081, A
16	53.6	4.7	1005	9 US-09-738-626-1392	Sequence 1392, Ap
17	53.6	4.7	3309400	9 US-09-738-626-1	Sequence 1, Appl1
18	53.2	4.7	418	10 US-09-960-352-4473	Sequence 4473, Ap
19	44.6	3.9	197	10 US-09-864-761-30128	Sequence 30128, A

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	21	43.2	3.8	1425	9	US-09-796-753-61	Sequence 61, Appl
	22	42.6	3.8	912	10	US-09-974-300-6823	Sequence 6823, Ap
C	23	39	3.4	592	9	US-09-902-563-5	Sequence 5, Appl1
C	24	39	3.4	5403	9	US-09-902-563-3	Sequence 3, Appl1
C	25	35.6	3.1	1642	9	US-10-153-668-71	Sequence 71, Appl
C	26	35.6	3.1	3427	9	US-10-153-668-73	Sequence 73, Appl
C	27	35.4	3.1	1149	10	US-09-880-107-3374	Sequence 3374, Ap
	28	35	3.1	427	10	US-09-983-965-357	Sequence 357, App
C	29	35	3.1	636	9	US-10-184-644-22	Sequence 22, Appl
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C	34	34.2	3.0	66804	10	US-09-740-041-3	Sequence 3, Appl1
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	39	33.8	3.0	1853	9	US-09-892-949-16	Sequence 16, Appl
	40	33.8	3.0	2402	9	US-09-892-949-15	Sequence 15, Appl1
	41	33.8	3.0	2529	9	US-09-892-949-45	Sequence 45, Appl
	42	33.8	3.0	2903	9	US-09-892-949-53	Sequence 53, Appl
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ALIGNMENTS

RESULT 1
US-10-091-628-1

Sequence 1, Application US/10091628

Patent No. US20020164627A1

GENERAL INFORMATION:

APPLICANT: Wilganowski, Nathaniel L.

APPLICANT: Nepomichy, Boris

APPLICANT: Burnett, Michael B.

APPLICANT: Hu, Yi

TITLE OF INVENTION: No. US20020164627A1 Human Transporter Proteins and Polynucleotides

FILE REFERENCE: LEX-0314-USA

CURRENT APPLICATION NUMBER: US/10/091,628

CURRENT FILING DATE: 2002-03-06

PRIOR APPLICATION NUMBER: US 60/275,009

PRIOR FILING DATE: 2001-03-12

PRIOR APPLICATION NUMBER: US 60/284,152

PRIOR FILING DATE: 2001-04-17

NUMBER OF SEQ ID NOS: 6

SOFTWARE: FastSeq for Windows Version 4.0

SEQ ID NO 1

LENGTH: 1134

TYPE: DNA

ORGANISM: Homo sapiens

US-10-091-628-1

Query Match 100.0%; Score 1134; DB 9; Length 1134;

Best Local Similarity 100.0%; Pred. No. 0;

Matches 1134; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Cy	1	ATGAGACCAATGTTTCACAGAGTCAGCCCTCCCAACGTTGAGAGGAGAGCTG	60
Db	1	ATGAGACCAATGTTTCACAGAGTCAGCCCTCCCAACGTTGAGAGGAGAGCTG	60
Cy	61	CCATGGAGCATGAGTGCATGGAACCTGAGCTCTTTTCACAGTGTCTCCACTGTG	120
Db	61	CCATGGAGCATGAGTGCATGGAACCTGAGCTCTTTTCACAGTGTCTCCACTGTG	120
Cy	121	ATGATGGAGGCTGCTCATGTTCTCTTGGAGTTCCTGGAATCCGGAAGCTGTGTCG	180
Db	121	ATGATGGAGGCTGCTCATGTTCTCTTGGAGTTCCTGGAATCCGGAAGCTGTGTCG	180

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QY 181 CACATCAGAGAGACCTGCGGAGATGCTGTGAGAGCTGCTGCGAGTTTGAGCTCATGCT 240
DB 181 CACATCAGAGAGAGACCTGCGGAGATGCTGTGAGAGCTGCTGCGAGTTTGAGCTCATGCT 240
QY 241 TTACAGCTTATCTGCGGAGATGCTGCTGAGAGCTGCTGAGAGCTGCTGAGAGCTGCT 300
DB 241 TTACAGCTTATCTGCGGAGATGCTGCTGAGAGCTGCTGAGAGCTGCTGAGAGCTGCT 300
QY 301 CTCATCATGAGGCTGCTGCGGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 360
DB 301 CTCATCATGAGGCTGCTGCGGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 360
QY 361 GAGATATGAGATCTGAGCATCAGATATGACACCTGTTCCACCGTGGCCGCTGGAGAG 420
DB 361 GAGATATGAGATCTGAGCATCAGATATGACACCTGTTCCACCGTGGCCGCTGGAGAG 420
QY 421 ATGCGCTGCTGAGTTATCTGACACCTGCTGAGAGCTGAGAGAGAGAGAGAGAGAGAGAG 480
DB 421 ATGCGCTGCTGAGTTATCTGACACCTGCTGAGAGCTGAGAGAGAGAGAGAGAGAGAGAG 480
QY 481 CCTTATCAGAGACATAGAGATTAACCTGTGTGCTGAGAGAGAGAGAGAGAGAGAGAGAG 540
DB 481 CCTTATCAGAGACATAGAGATTAACCTGTGTGCTGAGAGAGAGAGAGAGAGAGAGAGAG 540
QY 541 TATGTGAATTAAGATGAG 600
DB 541 TATGTGAATTAAGATGAG 600
QY 601 GAG 660
DB 601 GAG 660
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DB 661 AATTCAGAGATCAACCTTCTGAGCATCAGATGATGATGATGATGATGATGATGATGATG 720
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QY 841 GAGACATGAGAGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 900
DB 841 GAGACATGAGAGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 900
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RESULT 2

US-10-091-628-3
; Sequence 3, Application US/10091628
; Patent No. US20020164627A1
; GENERAL INFORMATION:
; APPLICANT: Walsanowski, Nathaniel L.
; APPLICANT: Nepomnichy, Boris
; APPLICANT: Burnett, Michael B.

APPLICANT: Hu, Yi
; TITLE OF INVENTION: No. US20020164627A1 Human Transporter Proteins and Polynucleoti-
; TITLE OF INVENTION: Same
; FILE REFERENCE: LEX-0314-USA
; CURRENT APPLICATION NUMBER: US/10/091,628
; PRIOR FILING DATE: 2002-03-06
; PRIOR APPLICATION NUMBER: US 60/275,009
; PRIOR FILING DATE: 2001-03-12
; PRIOR APPLICATION NUMBER: US 60/284,152
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO: 3
; LENGTH: 1600
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-091-628-3
Query Match 100.0%; Score 1134; DB 9; Length 1600;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 1134; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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DB 194 ATGAGAGCAATGTTTCAGAGAGCTGAGCTGAGCTGAGCTGAGCTGAGCTGAGCTGAGCTG 253
QY 61 CCAAGTGGAGCTGAG 120
DB 254 CCAAGTGGAGCTGAG 313
QY 121 ATGATGAGGCTGCTATGTTCTTGGAGAGTTCGAGAGAGAGAGAGAGAGAGAGAGAGAG 180
DB 314 ATGATGAGGCTGCTATGTTCTTGGAGAGTTCGAGAGAGAGAGAGAGAGAGAGAGAGAG 373
QY 181 CACATCAG 240
DB 374 CACATCAG 433
QY 241 TTACAGCTTATCTGCTGAGCATTTAGCTTTCTGAGAGAGAGAGAGAGAGAGAGAGAGAG 300
DB 434 TTACAGCTTATCTGCTGAGCATTTAGCTTTCTGAGAGAGAGAGAGAGAGAGAGAGAGAG 493
QY 301 CTCATCATGAGGCTGCTGCGGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 360
DB 494 CTCATCATGAGGCTGCTGCGGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 553
QY 361 GAGATATGAGATCTGAGCATCAGATATGACACCTGTTCCACCGTGGCCGCTGGAGAGAG 420
DB 554 GAGATATGAGATCTGAGCATCAGATATGACACCTGTTCCACCGTGGCCGCTGGAGAGAG 613
QY 421 ATGCACTCTGAGTTATCTTACACCTGCTGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 480
DB 614 ATGCACTCTGAGTTATCTTACACCTGCTGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 673
QY 481 CCTTATCAGAGATGAGATTAACCTGTGTGCTGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 540
DB 674 CCTTATCAGAGATGAGATTAACCTGTGTGCTGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 733
QY 541 TATGTGAATTAAGATGAG 600
DB 734 TATGTGAATTAAGATGAG 793
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DB 794 GATGAGGCTGCTTCTGAGTGTGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 853
QY 661 AATTCAGAGATCAACCTTCTGAGCATCAGATGATGATGATGATGATGATGATGATGATGAT 720
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QY 721 GATTTCTGCTGAGCATTTTACCACAGCTGTTGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 780
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Qy	781	GAACGTGAGCTCAGAAATATTCAGATGTCATCACCATGCTCCAGTATCTTCACTCT	840
Dd	974	GAACTGAGCTCAGAAATATTCAGATGTCATCACCATGCTCCAGTATCTTCACTCT	1033
Qy	841	GAGCATTGCTCAGATGTGAGTTCCACTGGGCTTATGCACTTCCAGCTGATGAT	900
Dd	1034	GAGCATTGCTCAGATGTGAGTTCCACTGGGCTTATGCACTTCCAGCTGATGAT	1099
Qy	901	GGATTCTTATTTGTTCAGCATATCAGACGTACAAGAGAGATTGAAGACAAACATGGA	960
Dd	1094	GGATTCTTATTTGTTCAGCATATCAGACGTACAAGAGAGATTGAAGACAAACATGGA	1153
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Dd	1154	AAAAAGAACTCAGGTTGCACAGAACTCTGCCATACAGAGAAATGCACTTCTTCCAGAGAG	1213
Qy	1021	ACCAATGCTCTTGTGAGGTGAATAGAAGGTGCATCACTCCGGGGCCACCAGGGCCA	1080
Dd	1214	ACCAATGCTCTTGTGAGGTGAATAGAAGGTGCATCACTCCGGGGCCACCAGGGCCA	1273
Qy	1081	ATGATTTGCCACAGGGCTCTCGAGCCAGTTGGCCCATCACTTCACTGGAATAG	1134
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: Sequence 626, Application US/09917800A
: Patent No. US20020119462A1
: GENERAL INFORMATION:
:   APPLICANT: Mendrick, Donna
:   APPLICANT: Porter, Mark
:   APPLICANT: Johnson, Kory
:   APPLICANT: Castle, Arthur
:   APPLICANT: Elashoff, Michael
:   APPLICANT: Gene Logic, Inc.
:   TITLE OF INVENTION: Molecular Toxicology Modeling
:   FILE REFERENCE: 44921-5038-US
:   CURRENT APPLICATION NUMBER: US/09/917,800A
:   CURRENT FILING DATE: 2001-07-31
:   PRIOR APPLICATION NUMBER: US 60/222,040
:   PRIOR FILING DATE: 2000-07-31
:   PRIOR APPLICATION NUMBER: US 60/222,880
:   PRIOR FILING DATE: 2000-11-02
:   PRIOR APPLICATION NUMBER: US 60/290,029
:   PRIOR FILING DATE: 2001-05-11
:   PRIOR APPLICATION NUMBER: US 60/290,645
:   PRIOR FILING DATE: 2001-05-15
:   PRIOR APPLICATION NUMBER: US 60/292,336
:   PRIOR FILING DATE: 2001-05-22
:   PRIOR APPLICATION NUMBER: US 60/295,798
:   PRIOR FILING DATE: 2001-06-06
:   PRIOR APPLICATION NUMBER: US 60/297,457
:   PRIOR FILING DATE: 2001-06-13
:   PRIOR APPLICATION NUMBER: US 60/298,884
:   PRIOR FILING DATE: 2001-06-19
:   PRIOR APPLICATION NUMBER: US 60/303,459
:   PRIOR FILING DATE: 2001-07-09
:   NUMBER OF SEQ ID NOS: 1740
:   SOFTWARE: PatentIn Ver. 2.1
:   SEQ ID NO 1626
:   LENGTH: 1663
:   TYPE: DNA
:   ORGANISM: Rattus norvegicus
:   FEATURE:
: OTHER INFORMATION: Genbank Accession No. US20020119462A1 NM_017047
: US-09-917-800A-1626

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				Gaps 2;

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Db	219	TAAATGCTCTTATCAATGCTCTCACTGGGCTGCACCATGGAATTCAGCAAGATCAAG	278
OY	179	CGCAGATCAGAGACCCTGGGGCAATGCTGTGGGAGTCTGTCCAGATTGGGCTATGC	238
Db	279	CTCACTTGTGGAAGCCCAAGGGGGATGATCGTTGCTTGATGGGCCCGATTGGGATCATGC	338
OY	229	CTTTTAGAGCTTATCTCTGGCATTAAGTTTCTCTGAAGCAGCCAAAGCATTTGCTG	298
Db	339	CCCTGCTGCTTTTCTCTCGCAAGATCTTTCACCTTGACAACATTTGAAGCTCTGGCA	398
OY	299	TTCTCATATGAGGCTGCTGCCCGGGGGGACCATCTCTAACTTTTACCTTCTGGGTTG	358
Db	399	TCTCATCTGTGTGCTGCTCTCCCGGGGGGAATTGTCCAACTCTTCAACCTCTGGCATGA	458
OY	359	ATGAGATATGAGATTCAGCATCAGTATGACAACCTGTTTCCACGTGGCCGCTTGGAA	418
Db	459	AGGGGAGATGAACCTCAGCATCTGATGAGCACCACTGCTCCAGCTTCAGTGCCTTGGCA	518
OY	419	TGATGCCACTCGCATTTATCTCTACAC--TGATCGTGAGTCTTGACAGATCTCA	475
Db	519	TGATGCCACTCTCTTAATACGTCTCAAGCAAGGACCTTAAGATGAGACCTTAAGACA	578
OY	476	CCATTCTTATCAGAACTAGGAATTAACCTTGTGTGTGCTGACATTCCTGTGGCCTTTG	535
Db	579	AGGTCCCTCAAGAGCATTAATGATATCACTAGTCAATGTTCTCATTTCTTGACCATAG	638
OY	536	GTTGTATATGATTAACAGATGGCCAAAACAATCCAAATCATTTCTCAAGATTGGGGCG	595
Db	639	GGATGTCTCTCAAGTCCAAAGGCCACACTATGTACCTTCAATCTTCAAGGGAGGATGA	698
OY	596	TTGTGTGGGGTCTCTCCCTTCTGTGTGTGTCAGTTGCTGTGTGTGGTCTGGCGAAAGAT	655
Db	699	TCATACCTCTCTCTCTGTGTGTGTGTCAGCCCTCTCTGTATCAATGTGTGGCAACA	758
OY	656	CTTGAATTCAGACATCAC-----CCTTGTGACCATCAGTTTCATCTTTCTTTGATTG	709
Db	759	GCATCATGTGTGTCATGACACCACTAATGCTAATCTCTCCCGATGACCTTCTCTG	818
OY	710	GGCATGTCAAGGGTTTCTGTGTGCACTTTTACCAACCAAGTCTTGGCAAAGTGCAGA	769
Db	819	GCTTCTGATGGGTAACTCTCTGTGCTCTTCTTCAATCAATCCAACTGAGACGA	878
OY	770	CAATTCTTGAAGATGAGCTCAGAAATATTCAGATGTGCATCCATGCTCCAGTTAT	829
Db	879	CCATGAGATGGAACAGGATTCCAAACATTCACACTGTGTCTACATCTCTCAATGTGA	938
OY	830	CTTTCACGTGAGACCTTGTGTCCAGATGTAGTTTCCCACTGAGCTTAGACCTTCC	889
Db	939	CCCTTCCCCCTGAAATCAATGGGCCACTTTTCTTTCTCTCTCTCTCAATGATTTCC	998
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Db	999	AGCTTGCAAGAGACTTTCAT	1020

RESULT 4

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US-09-880-107-2176
Sequence 2176, Application US/09880107
Patent No. US20020142981A1
GENERAL INFORMATION:
APPLICANT: Horne, Darci T.
APPLICANT: Vockley, Joseph G.
APPLICANT: Scherf, Uwe
APPLICANT: Gene Logic, Inc.
TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer
FILE REFERENCE: 44921-5028-WO
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR FILING DATE: 2000-06-14
PRIOR APPLICATION NUMBER: US 60/237,054

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PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2176
LENGTH: 1580
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
OTHER INFORMATION: Genbank Accession No. US20020142981A1 L21893
US-09-880-107-2176

Query Match 15.3%; Score 173.6; DB 10; Length 1580;
Best Local Similarity 51.9%; Pred. No. 1e-46;
Matches 445; Conservative 0; Mismatches 404; Indels 9; Gaps 2;

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179 CGACATCAGGAGACCTGGGGCATGCTGTGGACTGCTTCCAGTTTGGGCTCATGC 238
240 CTCACCTTAAGAGCCCTAAAGGCTGGCCATCCCTGGTGGCAGATATGGCATATGC 299
239 CTTTACAGCTTATCTCTGGCCATTAGCTTTCTCTGAAGCCAGTCCAGCTATTGTG 298
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299 TTCTCATATGAGGCTGCTGCGCGGGGAGCACTCTCTCAATTTTCACTTCTGGGTTG 358
360 TCTGTGCTGTGCTGCTGCTGCTGAGGAGAACTGTCAATGCTTCAATGCTGGCATGA 419
359 ATGGAATATGATCTCAGATCATATGAAACCTGTTCCACCGTGGCCGCTTGGAA 418
420 AGGGGAGCAATGAACCTCAGCATGTATGATGACCACTGCTCCACTTGTGCTTGGCA 479
419 TGATGACACTGCTGCTTATCTCTACACC---TGATCGAGATCTTCAAGCAATCTCA 475
480 TGAATGCTCTCTCTCTGATCATTTACTCCAGGGGATCTATATGAGGAGCTTGAAGACA 539
476 CCAATTCCTTATAGAAATGAAATTAACCTTGTGCTGACATCTCTGTCGCTTTG 535
540 AGGTGCCCTATAAAGCATGTATATCACTGTGCTGCTGCTCAATCTTCTGACCATAG 599
536 GTGCTATGATGATTAAGATGAGGCAAAACATCCAAATCATTTCTCAAGATTGGGCGG 595
600 GGAATGCTCTCAATATCAACGCGCAATATGCGCTATGTCATCAAGGAGGATGA 659
596 TTGTTGATGGGCTCTCTCTGATGATGAGTGTGCTGCTGCTGCGCAAGAT 655
660 TCAATATCTCTTGTGAGTGTGCGCTGACATTTCTCTGCAATCAATGTGGGAAAGA 719
656 CTTGAATTCAGATCAC-----CCTTCAAGCATCAATTTCTTCTTCTTGAATG 709
720 GCATCATGTTTGCATGACACCACTCTTGAATGCCACCTCCCTCGATGCCCTTATG 779
710 GCCATGTCAGGGTTTGTGCTGCGCACTTTTACCAACAGTCTTGGCAAGGTGACAGA 769
780 GCTTCTGTGGGTATGTTTCTCTGCTCTCTTCTTCTCAATGAGCGGTGACAGCA 839
770 CAATTCCTTGAATCGAGCTCAGAAATTCAGATGTCATCAGATGCTCAGATTAT 829
840 CTGTACGATGAGATGAGATGCAAAATGTCCAACTGTCTTCCACCACTCTCAATGTGG 899
830 CTTTCACTGCTGAGCACTTGTGTCAGATGTTGATTTCCACATGGCCTTATGACTCTCC 889
900 CTTTCCACCTGAAGTATGAGCACTTTTCTTCTTCTTCCCTCCCTCAATCATATTTTCC 959
890 AGTGAATAGTGAATTTCTTATTTGTGACATATAGACATCAAGAGATGTTGAAGA 949
960 AGCTTGAGAAAGGGCTTCTCTCATTTGTCATATTTTGGTCTATGAGAAATTCAGACTC 1019
950 ACAACATGGAAGAAAGA 967

DB 1020 CCAAGATRAAAACAAAA 1037

RESULT 5
US-09-864-761-31375/c
Sequence 31375, Application US/09864761
Patent No. US20020048763A1
GENERAL INFORMATION:
APPLICANT: Penn, Sharon G.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
FILE REFERENCE: Aecm1ca-X-1
CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00662
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR FILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: US 09/774,203
NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
SEQ ID NO 31375
LENGTH: 360
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
OTHER INFORMATION: MAP TO AL157789.1
OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 2.2
OTHER INFORMATION: NT HIT: g11435250, EVALUE 0.00e+00
OTHER INFORMATION: SWISSPROT HIT: Q14973, EVALUE 7.00e-64
OTHER INFORMATION: EST_HUMAN HIT: W01479.1, EVALUE 0.00e+00
US-09-864-761-31375

Query Match 7.0%; Score 79.8; DB 10; Length 360;
Best Local Similarity 56.8%; Pred. No. 6.7e-16;
Matches 147; Conservative 0; Mismatches 112; Indels 0; Gaps 0;

119 TGAATGAGGAGCTGCTCATGTTCTCTTGGAGTTCCTGGAGATCCGGAAGCTGTGT 178

Db 41 ACATACAGAAATGTTATGAAAAATGATG 11

RESULT 8

US-09-960-352-2253
Sequence 2253, Application US/09960352
Patent No. US20020137139A1
GENERAL INFORMATION:
APPLICANT: Warren, Wesley C.
APPLICANT: Tao, Nengbing
APPLICANT: Byatt, John C.
APPLICANT: Mathialagan, Nagappan
TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
FILE OF INVENTION: MUSCLE AND FAT DEPOSITION
FILE REFERENCE: 16511.006/37-21(10298)C
CURRENT APPLICATION NUMBER: US/09/960,352
CURRENT FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 15112
SEQ ID NO 2253
TYPE: DNA
ORGANISM: Bos taurus
FEATURE:
NAME/KEY: unsure
LOCATION: (390)
OTHER INFORMATION: unsure at all n locations
OTHER INFORMATION: Clone ID: 10-LIB34-014-01-E1-C5
US-09-960-352-2253

Query Match Best Local Similarity 6.5%; Score 74; DB 10; Length 401;

Matches 122; Conservative 0; Mismatches 81; Indels 0; Gaps 0;

QY 119 TGATGATGAGGGGCTGCTGATGTTCTCTTGGGATGTTCCGAGGATCCGGAAGCTGTGT 178
Db 199 TCATGCTGTTACCATGATGCTCTCGCTGGGTGACCATGAGTTTCAGCAGATCAAG 258
QY 179 CGCATCATGAGAGACCCCTGGGGCATGCTGTGGAGCTGCTGCCAGTTTGGGCTCATGC 238
Db 259 CGCATCTTGGAGAGCCCAAGGGGCTGGCGTGGTGGGCGCAGTTTGGGCTCATGC 318
QY 239 CTTTACAGCTTATCTCTCTGCGCATTAAGCTTTCTCTGAAGCCAGTCCAGCTATGCTG 298
Db 319 CCTCCTGCTGCTTGGAGTGGGCAAGTTCTTCCAGCTGAATTAAGTTGAGGCCCTTACCA 378
QY 299 TTCTCATCATGAGGCTGTGCGCCG 321
Db 379 TCCTGATCTGCNCTGCTCTCACCG 401

RESULT 9

US-09-738-626-2554
Sequence 2554, Application US/09738626
Publication No. US20020197605A1
GENERAL INFORMATION:
APPLICANT: NAKAGAWA, SATOSHI
APPLICANT: MIZOGUCHI, HIROSHI
APPLICANT: ANDO, SEIKO
APPLICANT: HAYASHI, MIKIRO
APPLICANT: OCHIAI, KEIKO
APPLICANT: YOKOI, HARUHIKO
APPLICANT: TATEISHI, NAKO
APPLICANT: SENOH, AKIHIRO
APPLICANT: IKEDA, MASATO
APPLICANT: OZAKI, AKIO
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES
FILE REFERENCE: 249-125
CURRENT APPLICATION NUMBER: US/09/738,626
CURRENT FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: JP 99/377484
PRIOR FILING DATE: 1999-12-16
PRIOR APPLICATION NUMBER: JP 00/159162

PRIOR FILING DATE: 2000-04-07
PRIOR APPLICATION NUMBER: JP 00/280988
PRIOR FILING DATE: 2000-08-03
NUMBER OF SEQ ID NOS: 7059
SOFTWARE: PatentIn ver. 3.0
SEQ ID NO 2554
LENGTH: 972
TYPE: DNA
ORGANISM: Corynebacterium glutamicum
US-09-738-626-2554

Query Match Best Local Similarity 6.0%; Score 67.6; DB 9; Length 972;

Matches 157; Conservative 0; Mismatches 149; Indels 0; Gaps 0;

QY 129 GCTGCTCATGTTCTCTTGGGATGTTCCGAGATCCGGAAGCTGTGTCACATGAG 188
Db 141 GATCATCATGTTTACCATGAGTTTACCTTGACGGTGGCCGATTTTCAGATGTCCTTA 200
QY 189 GAGACCTGGGACATGCTGTGGGACCTGCTGCGCATTTGGGCTCATGCTTTTACAGC 248
Db 201 ACGTCCACTGCTTATCTTGAATCGGTGTAGTACGCGCATTTGTCATCATGCGCATTCCTGGC 260
QY 249 TTAATCTCCGACCATTAAGCTTTTCTGTAAGCCAGTCCAGCTATTGCTGTCTCATAT 308
Db 261 GATCGGTGTGGAAATGTTCAACCTCAACCCAGCAGCTGCGGTGGCTTCTCATGCT 320
QY 309 GGGCTGCTGCCCCGGGGGACCATCTCTACATTTTCACTTCTGGGTTGATGAGATAT 368
Db 321 GGGATCCGTTCCGGGTGGGACCTCTTCCAGTATGATGGTTTCTGGCCGAGAGATGT 380
QY 369 GGATTCAGCATGATGATGACCACTGTTCACCGTGGCCGCTGGGAGATATGCCACT 428-
Db 381 CGCGCTATCGGTCAACCATGACCTCTGTGTCCACCATTTTCCCAATCATGACCGCTTT 440
QY 429 CTGCAAT 434
Db 441 CCTCAT 446

RESULT 10

US-09-738-626-1/c
Sequence 1, Application US/09738626
Publication No. US20020197605A1
GENERAL INFORMATION:
APPLICANT: NAKAGAWA, SATOSHI
APPLICANT: MIZOGUCHI, HIROSHI
APPLICANT: ANDO, SEIKO
APPLICANT: HAYASHI, MIKIRO
APPLICANT: OCHIAI, KEIKO
APPLICANT: YOKOI, HARUHIKO
APPLICANT: TATEISHI, NAKO
APPLICANT: SENOH, AKIHIRO
APPLICANT: IKEDA, MASATO
APPLICANT: OZAKI, AKIO
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES
FILE REFERENCE: 249-125
CURRENT APPLICATION NUMBER: US/09/738,626
CURRENT FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: JP 99/377484
PRIOR FILING DATE: 1999-12-16
PRIOR APPLICATION NUMBER: JP 00/159162
PRIOR FILING DATE: 2000-04-07
PRIOR APPLICATION NUMBER: JP 00/280988
PRIOR FILING DATE: 2000-08-03
NUMBER OF SEQ ID NOS: 7059
SOFTWARE: PatentIn ver. 3.0
SEQ ID NO 1
LENGTH: 3309400
TYPE: DNA
ORGANISM: Corynebacterium glutamicum
US-09-738-626-1


```
RESULT 13
US-10-091-628-6
; Sequence 6, Application US/10091628
; Patent No. US20020164627A1
; GENERAL INFORMATION:
; APPLICANT: Wiganowski, Nathaniel L.
; APPLICANT: Nepomichy, Boris
; APPLICANT: Burnett, Michael B.
; APPLICANT: Hu, Yi
; TITLE OF INVENTION: No. US20020164627A1el Human Transporter Proteins and Polynucleoti
; TITLE OF INVENTION: Same
; FILE REFERENCE: Lex-0314-USA
; CURRENT APPLICATION NUMBER: US/10/091,628
; PRIOR FILING DATE: 2002-03-06
; PRIOR APPLICATION NUMBER: US 60/275,009
; PRIOR FILING DATE: 2001-03-12
; PRIOR APPLICATION NUMBER: US 60/284,152
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 6
; LENGTH: 1777
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-091-628-6

Query Match
Best Local Similarity 5.5%; Score 62.4; DB 9; Length 1777;
Matches 207; Conservative 0; Mismatches 241; Indels 0; Gaps 0;

QY 131 TGTCTATGTTCTTTGGATGTTCCGTGAGATCCGGAAGCTGTGTCGACATCAGGA 190
DB 756 TGAATATAGTGATGTTGGTTGTAAGATGAATTACAGCTGTTTCAACAGATGGAAGA 815
QY 191 GACCTGGGGCATGCTGTGGGACTGCTGCCAGTTTGGGCTCATGCTTTTACAGCTT 250
DB 816 GACCTTGGCAGTAATCTTTGGGGCAGTTACACAGTTTTTTCGATGCCATTTTGGGGGT 875
QY 251 ATCTCTGGCANTTACCTTTTCTGAGGCACTTCCAGCACTTCCAGCTTCTTCATCATG 310
DB 876 TTTCTTTGCTCAGATTGTCGATTCCTGAGCGGAGCTTTTGGAGTTGTAATACCT 935
QY 311 GCTGTCGCCCGGGGCGACCATCTTCAATTTTCACTTCTGGGTTGATGAGATATG 370
DB 936 GCACTGCGCCAGGAGGGGGGTATCTCTTTGCTGCTTGAATGAGATTTCA 995
QY 371 ATCTCAGCATCAGTATGACACTGTTCCACCGTGGCCGCTGGGAATGATGCCACTCT 430
DB 996 CATGGCCATTTGATGACTTGCACATCAATATTGGCTGATCATGATGCCCTGTCA 1055
QY 431 GCATTATCTCTACACTGCTGCTGAGTCTTCAAGAGATCCACATTCCTTATCAGA 490
DB 1056 ATTCTTATATATACAGTATGAGATTTAGGCTTGCAGTATTCATATTCGTTTCTA 1115
QY 491 ACATAGAAATTAACCTTGTGTGCTGACCAATTCCTGTGGCTTGTGTATGTAAT 550
DB 1116 AATTTGTGCAACACTCTTTTCACTTGTGTGCGAGATCAATTGAAATAGTATCAAGC 1175
QY 551 ACAGATGGCCAAACATCCAAATCAT 578
DB 1176 ATAGATTAACCTGAAAAAGCAAGCTTCTT 1203

RESULT 14
US-09-833-381-318
; Sequence 318, Application US/09833381
; Patent No. US20020132090A1
; GENERAL INFORMATION:
; APPLICANT: Robison, Keith E.
; TITLE OF INVENTION: No. US20020132090A1el Nucleic Acid and Protein Homologs
; FILE REFERENCE: 5800-119
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; CURRENT APPLICATION NUMBER: US/09/833,381
; CURRENT FILING DATE: 2001-04-11
; PRIOR APPLICATION NUMBER: 09/516,448
; PRIOR FILING DATE: 2000-02-29
; NUMBER OF SEQ ID NOS: 2050
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 318
; LENGTH: 374
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(374)
; OTHER INFORMATION: n = A,T,C or G
US-09-833-381-318

Query Match
Best Local Similarity 5.2%; Score 58.6; DB 10; Length 374;
Matches 141; Conservative 0; Mismatches 100; Indels 3; Gaps 2;

QY 80 ATGAAACCTGAGCTCGTTTTCACAGTGTCCACTGTGATGATGGGCTCTCATGT 139
DB 131 ATGCAATTCCTCAATACAGTATGAGCACTGTGCTCATCCTTTAGCATGTGATGT 190
QY 140 TCTCTTGGAGATGTTCCGTGAGATCCGGAAGCTGTGTCGACATCAGAGACCTTGG 199
DB 191 TTTCTATGGGGGTGCAATGTGAGAGTCACAAAGTTCTTGAACATTAAGAGACATGG 250
QY 200 GCATTGCTGTGGAGCATGCTGCTGCGCAGTTTGGGCTCATGCTTT--TACAGTTATCTCT 257
DB 251 GTATCTGTGGGCTTCTCTGTCTGTCAAGTTTGAATATACCTCTCCAAAGCTTTATAC 310
QY 258 GGCATTAGCTTTTCTCTGAAAGCCAGTCCAAAGCTATTGCT--GTTCTCATATGAGCTCT 316
DB 311 TGTCTGTGGCCCTGTGATCTTCTCTGACAGCTGTGATGTGCTAATTATATGGGTGCT 370
QY 317 GCCC 320
DB 371 GCCC 374

RESULT 15
US-09-960-352-10081
; Sequence 10081, Application US/09960352
; Patent No. US20020137139A1
; GENERAL INFORMATION:
; APPLICANT: Warren, Wesley C.
; APPLICANT: Tao, Nengping
; APPLICANT: Byatt, John C.
; APPLICANT: Mathialagan, Nagappan
; TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
; FILE REFERENCE: 16511.006/37-21(10298)C
; CURRENT APPLICATION NUMBER: US/09/960,352
; CURRENT FILING DATE: 2001-09-24
; NUMBER OF SEQ ID NOS: 15112
; SEQ ID NO 10081
; LENGTH: 407
; TYPE: DNA
; ORGANISM: Bos taurus
; OTHER INFORMATION: Clone ID: 43-LIB34-043-Q1-E1-C4
US-09-960-352-10081

Query Match
Best Local Similarity 4.8%; Score 54.2; DB 10; Length 407;
Matches 120; Conservative 0; Mismatches 88; Indels 3; Gaps 1;

QY 359 ATGAGATATGATGATTCAGCATCAGTATGACAACTGTTCCACCGTGGCCCTGGGAA 418
DB 10 AGGGGAGACATGAACCTGAGCATGAGTATGACCACTGTCTCCACCTTTCGCGCTGGGCA 69
QY 419 TGATGCCACTGTCATTTATCTCTACACTGCTGAGTCTTCTAG---CAGATTCGA 475
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Db 70 TGATGCCCCCTCCTCCTGTACTTACTCCAGGGGCATCTATGATGGGTCCCTGAAGACA 129
 Oy 476 CCATTCTTATCAGAACATAGGAATTACCCCTTGTGTGCTGACCATTCCTGTGGCCTTG 535
 Db 130 AGGTGCGGTACGGCGGCATCATGATATCATCTGATCTCTGATTCCTCATCCCTTGACCATAG 189
 Oy 536 GTGTCTATGTGATTAACAGATGGCCAAACA 566
 Db 190 GCATCATCCTCAATCCAAAGGCCCAATA 220

Search completed: June 9, 2003, 06:34:43
 Job time : 430 secs

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